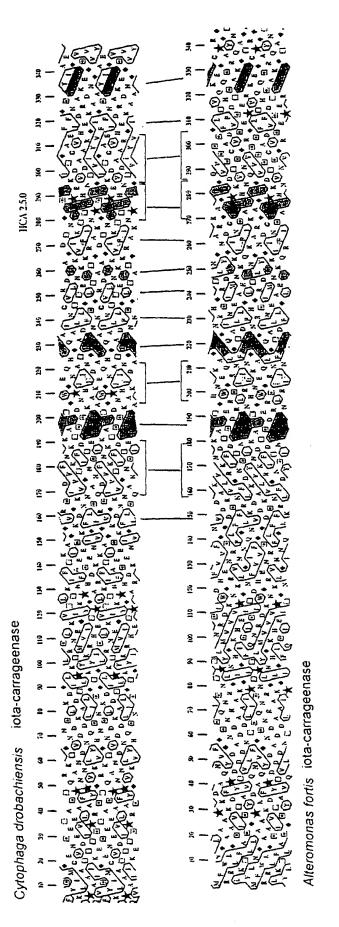
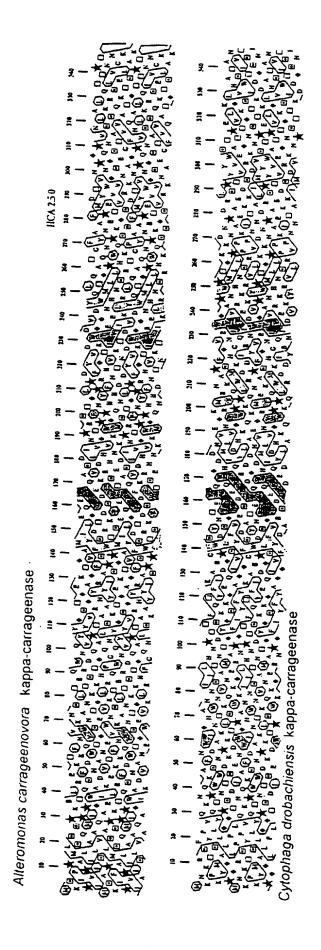
	↓ ↓ ↓	
27	AVSPKTYKDADFYVAPTQQDVNYDLVDDFGANGNDTSD	64
28	DISEISEVPIELRAAASSFYTPPGQNVRANKKNLVIDYGVNHNDQND ↓	74
65	DSNALQRAINAISRKPNOGILLIPNGIYHFIGIQMKSNVHIRVESDVIIK	114
75	DSSKLNLAIKDLSDIGGILITLPKGKYYLIKIRMRSNVHLEIEKGIVIY	122
115	PIWN GDGKNHRLFEVGVNNIVRNFSFQGLGNGFLVDFKDSRDKNLAV	161
123	PIKGLIPAKNHRIFDFASKIEEKIENASIVGKGGKFIVDLRGNSSKNQIV	172
162	FKLGDVRNYKISNFTIDDNKTIFASILVDVTERNGRLHWSRNGIIERIKQ	211
173	ADVGNVINFKISNFTIKDEKTIFASILVSFTDKAGN_AWPHKGIIENIDQ	221
212	NNALFGYGLIQIYGADNILFRNLHSEGGIALRMEITNILMKNYKQGIRN	261
222	ANAHIGYGLIQAYAALNILFNNLSCIGGVTLRLETENLAMKTAKKGGVRD	271
262	IFADNIRCSKGLAAVMFGPHFMKNGDVQVINVSSVSCGSAVRSDSGFVEL	311
272	IFATKIKNINGLTPVMFSPHFMENGKVTIDDVTAIGCAYAVRVEHGFIEI	321
312	FSPIDEVHIRQSWKQAVESKLGRCCAQIPYARGNOGIRWAARVT OKD	358
322	FDKGNRASA_DAFKNYIEGILGAGSVEVVYKRNNGRT_WAARIANDFNEA	369
359	ACLDKAKLEYGIEPGSFGIVKVFDVIARF_GYNADLKQDQLDYFSISNPM	407
370	AYNHSNPAVSGIKPCKFATSKVINVKATYKGICAKLKQAFLSYLPCSER_	418
408	CKRVCLPTKEQWSKQGQIYIGPSLAAVID_TTPETSKYDYDVKTFNVKRI	457
419		462
457	NFPVNSHKTIDINIESSRVCNYY_GMSECSSSRWER	491
463	GFPNNYVLNVKYNT_PKVCNQNLG_SITSCN	491

FIG.1



1.0 H

1	MKKPNFYGKMGRTALSSLFYLFFLGLVYGQQPTKTSNPNDQWTIKWSASDEFN_KNDPDW	59
1	MKPISIVAFPIPAISMLLLSAVSQAASM_QPPIAK_PGETWILQAKRSDEFNVK_DAT_	55
60	AKWIK TGNLPNTSAWKWN NOKNVKISNGIAELTM RHNANNTPPDGGT YF	108
56		109
109	_TSGIFKSYQKFTYGYFEAKIQGADIGEGVCPSFWLYSDFDYSVAN_GETVYSEIDVVEL	166
110	YTSGVAKSRATGNYGYYEARIKGASTFPGVSPAFWMYSTIDRSLTKEGDVQYSEIDVVEL	169
167	QQFDWY_EGHQDDIYDMDLNLHAVVKENGQGVWKRPKMYPQEQLNKWRAM_DPSKDFHIY	224
170	TOKSAVRES DH DLH NI VVK NGKPTWMRPGSFPQTNHNGYHLPFDPRNDFHTY	221
225	GCEVNONEIIWYVDGVE_VARKPNKYWHRPMNVTLSLGLRKPFVKFFDNKNNAINPETDA	283
222	GVNVTKDKITWYVDG_EIVGEKDNLYWHRQMNLTLSQGLRAPHTQWKCNQFYPSAN_	276
	K_AREKLSDIPTSMYVDYVRVWEKSAGNTTNPPTSEVGTLKTKGSKLVIDHWDASTGTIS	342
277	KSA_EGFPTSMEVDYVRTWVKVGNNNSAPGEGQSCPNTFVAVNSVQLSAAKQTLRKG	332
343	AVSNNTKTGQYAGSVNNASIAQIVTLKANTSYKVSAFGKASSPGTSAYLGISKASNNELI	402
333	QSTTLESTVLPNCATNKKVIYSSSNKNVATVNSAGVV_KAKNKGTATITVKTKNKGKIDKL	392
403	SNFEFKTTSYSKGEIEIRTGNVQESYRIWYWSSGQAYCDDFNLVEINSGASQLNENETET	462
393	TIAVN	397
463	ALEKGIHIYPNPYKNGPLTIDFGKPFSGEVQITGLNGRTFLRRNVVDQTSVQLLESKSKF	522
523	KSGLYIVKISGPDGEVSKKILVE	545



F16.4